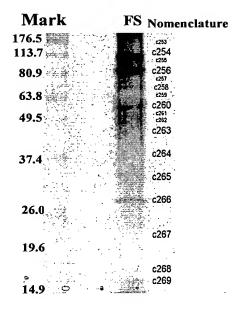
Figure 1. Isolation of human Fibrous Sheath



Figure 2. SDS-PAGE of human fibrous sheath proteins



## Figure 3. Microsequencing of Fibrous Sheath Proteins

Each band of fibrous sheath protein was microsequenced by mass spectrometry. The sequence result was summarized in table 1. The band C265 was identified as an unknown protein (DKFZp434N1235). Peptides microsequenced from the C265 band are indicated by bold.

## Microsequencing of C265 by Mass Spectrometry

mhrepakka ekr**lfdassf gkdllaggva aavsk**tavap iervklllqv qasskqispe arykgmvdel vripreqgff sfwrgnlanv iryfptqaln fafkdkykql fmsgvnkekq fwrwflanla sggaagatsl cvvypldfar trlgvdigkg peerqfkglg dcimkiaksd giaglyqgfg vsvqgiivyr **asyfgaydtv k**gllpkpkkt pflvsffiaq vvttcsgils ypfdtvrrrm mmqsgeakrq yk**gtldcfvk iyqhegissf fr**gafsnvlr gtggalvlvl ydkikeffhi diggr

## **Mouse Orthologues of SFEC**

msnesskkqs skkalfdpvs fskdllaggv aaavskttva piervklllq vqasskqisp earykgmldc lvripreqgf lsywrgnlan viryfptqal nfafkdkyke lfmsgvnkek qfwrwflanl asggaagats lcvvypldfa rtrlgvdigk gpeqrqftgl gdcimkiaks dgliglyqgf gvsvqgiivy rasyfgaydt vkgllpkpke tpflvsfiia qivttcsgil sypfdtvrrr mmmqsgesdr qykgtidcfl kiyrhegvpa ffrgafsnil rgtggalvlv lydkikefln idvggsssgd

Figure 4. SFEC is a testis specific Protein- Northern Analysis

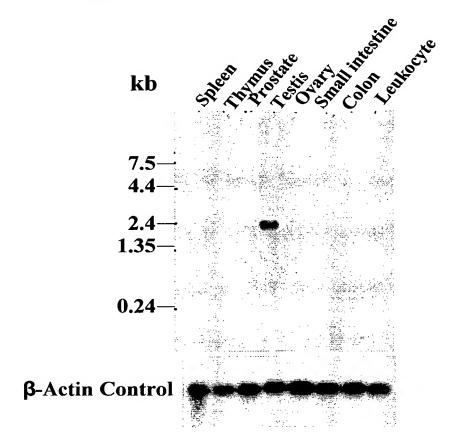


Figure 5- Dot blot analysis (upper panel) and human MTE array (lower panel) demonstrating testis specific expression of SFEC

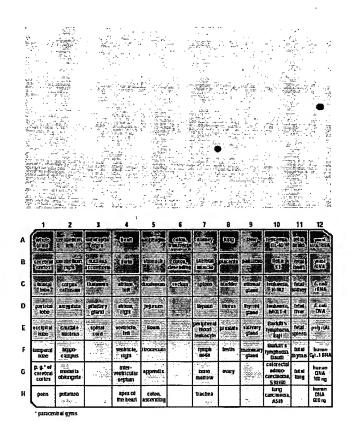
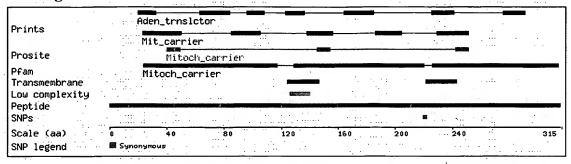


Figure 6- Functional Domains of SFEC



Domain type	Accession number Description	Start End
prints prints	PR00926         Mit_carrier           PR00926         Mit_carrier	23 36 50
prints	PR00926 Mit_carrier	B5 105
prints	PR00926         Mit_carrier           PR00926         Mit carrier	138
prints prints	PR00926         Mit_carrier           PR00926         Mit_carrier	229 251
prints prints	PR00927         Aden_tmslctor           PR00927         Aden_tmslctor	20 32 63 84
prints	PR00927 Aden_trnslctor	96 2 108
prints prints	PR00927         Aden_trnslctor           PR00927         Aden_trnslctor	123 136 136 164 185 185 185 185 185 185 185 185 185 185
prints	PR00927 Aden_trnslctor	225
prints Pfam	PR00927 Aden_trnslctor PF00153 Mitoch carrier	···· 275 · · · · · · · · · · · · · · · · · · ·
Pfam	PF00153 Mitoch_carrier	129 220
Pfam prosite	PF00153 Mitoch_carrier PS00215 Mitoch_carrier	226 314 40 49
prosite prosite	<u>PS00215</u> Mitoch_carrier <u>PS00215</u> Mitoch_carrier	145 154 242 251

Domain type		End
Transmembrane	124	146
Transmembrane	221 y with X	243
Low complexity	126	140

Figure 7. Alignment of Amino Acid Sequences of SFEC with other human proteins having a similar domains

CLUSTAL W (1.74) multiple sequence alignment unk|VIRT5947|Blast\_submission MHREPAKKKAEKRLFDASSFGKDLLAGGVAAAVSKTAVAPIERVKLLLQV MHREPAKKKAEKRLFDASSFGKDLLAGGVAAAVSKTAVAPIERVKLLLQV tr109H0C2 sp|P12235|ADT1 HUMAN -----MGDHAUSFLKDFLAGGVAAAVSKTAVAPIERVKLLLQV -----MTEQAISFAKDFLAGGIAAAISKTAVAPIERVKLLLQV sp|P12236|ADT3 HUMAN -----HTDAAVSFAKDFLAGGVAAAISKTAVAPIERVKLLLQV sp|PO5141|ADT2\_HUMAN \* \*\* \*\*: \*\*\*: \*\*\*: \*\*\*\*\*\*\*\*\* unk|VIRT5947|Blast submission QASSKQISPEARYKGMVDCLVRIPREQGFFSFWRGNLANVIRYFPTQALN OASSKOISPEARYKGMVDCLVRIPREOGFFSFWRGNLANVIRYFPTOALN tr|Q9H0C2 sp|P12235|ADT1 HUMAN QHASKQISAEKQYKGIIDCVVRIPKEQGFLSFWRGNLANVIRYFPTQALN sp|P12236|ADT3 HUMAN QHASKQIAADKQYKGIVDCIVRIPKEQGVLSFWRGNLANVIRYFPTQALN sp|PO5141|ADT2 HUMAN QHASKQITADKQYKGIIDCVVRIPKEQGVLSFWRGNLANVIRYFPTQALN \* :\*\*\*\*:.: :\*\*\*:\*\*:\*\*\*:\*\*\*.:\*\*\*\*\*\*\*\*\*\* unk[VIRT5947|Blast submission FAFKDKYKQLFMSGVNKEKQFURUFLANLASGGAAGATSLCVVYPLDFAR FAFKDKYKQLFMSGVNKEKQFURUFLANLASGGAAGATSLCVVYPLDFAR tr|Q9H0C2 sp|P12235|ADT1 HUMAN FAFKDKYKQLFLGGVDRHKQFURYFAGNLASGGAAGATSLCFVYPLDFAR sp|P12236|ADT3\_HUMAN  ${\tt FAFKDKYKQIFLGGVDKHTQFURYFAGNLASGGAAGATSLCFVYPLDFAR}$  ${\tt FAFKDKYKQIFLGGVDKRTQFURYFAGNLASGGAAGATSLCFVYPLDFAR}$ sp|PO5141|ADT2\_HUMAN \* TRLGVDIGKGPEERQFKGLGDCIMKIAKSDGIAGLYQGFGVSVQGIIVYR unk|VIRT5947|Blast\_submission TRLGVDIGKGPEERQFKGLGDCIMKIAKSDGIAGLYQGFGVSVQGIIVYR tr|Q9H0C2 sp| P12235| ADT1\_HUMAN TRLAADVGKGAAOREFHGLGDCIIKIFKSDGLRGLYOGFNVSVOGIIIYR sp|P12236|ADT3\_HUMAN TRLAADVGKSGTEREFRGLGDCLVKITKSDGIRGLYQGFSVSVQGIIIYR sp|PO5141|ADT2 HUMAN TRLAADVGKAGAEREFRGLGDCLVKIYKSDGIKGLYQGFNVSVQGIIIYR unk|VIRT5947|Blast\_submission ASYFGAYDTVKGLLPKPKKTPFLVSFFIAOVVTTCSGILSYPFDTVRRRM **ASYFGAYDTVKGLLPKPKKTPFLVSFFIAQVVTTCSGILSYPFDTVRRRM** tr | Q9H0C2 sp|P12235|ADT1 HUMAN **AAYFGVYDTAKGNLPDPKNVHIFVSUMIAQSVTAVAGLVSYPFDTVRRRM** sp | P12236 | ADT3 HUMAN **AAYFGVYDTAKGMLPDPKNTHIVVSUMIAQTVTAVAGVVSYPFDTVRRRM** sp|PO5141|ADT2\_HUMAN AAYFGIYDTAKGNLPDPKNTHIVISUMIAQTVTAVAGLTSYPFDTVRRRM \*:\*\*\* \*\*\*.\*\*:\*\*,\*\*;. :.:\*::\*\*\* \*\*: :\*: \*\*\*\*\*\* MMQSGE--AKRQYKGTLDCFVKIYQHEGISSFFRGAFSNVLRGTGGALVL unk|VIRT5947|Blast\_submission  ${\tt MMQSGE--AKRQYKGTLDCFVKIYQHEGISSFFRGAFSNVLRGTGGALVL}$ tr|Q9H0C2 sp | P12235 | ADT1\_HUMAN MMOSGRKGADIMYTGTVDCWRKIAKDEGAKAFFKGAWSNVLRGMGGAFVL MMQSGRKGADIMYTGTVDCWRKIFRDEGGKAFFKGAWSNVLRGMGGAFVL spiP12236|ADT3 HUMAN sp|PO5141|ADT2 HUMAN MMQSGRKGTD IMYTGTLDCWRKIARDEGGKAFFKGAUSNVLRGMGGAFVL \*.\*\*:\*\*: \*\* :.\*\* .:\*\*:\*\*:\*\*\*\* \*\*\*:\*\* VLYDKIKEFFHIDIGGR unk|VIRT5947|Blast\_submission VLYDKIKEFFHIDIGGR tr/Q9H0C2 VLYDEIKKYV---sp | P12235 | ADT1 HUMAN sp | P12236 | ADT3\_HUMAN VLYDELKKVI-----VLYDEIKKYT----sp|PO5141|ADT2 HUMAN

> Heart (ADT 1): identity (69%) similarity (79%) Liver (ADT 3): identity (67%), similarity (80%) Fibroblast (ADT 2): identity (67%), similarity (79%)

Figure 8. Human Fibrous Sheath Peptides involved in Energy Production

Gene/Gene	Peptides	Gene	Tissue
Symbol		Locus	Distribution
Aldolase A,	1)GILAADESTGSIAK	16q22-q24	Ubiquitously
Fructose-	2)IGEHTPSALAIMENANVLAR		expressed
bisphosphate	3)GVVPLAGTNGETTTQGLDGLSER		
ALDOA	4)FSHEEIAMATVTALR		
	5)IGEHTPSALAIMENANVLAR		
Pyruvate Kinase	1)NTGIICTIGPASR	15q22CM	Ubiquitously
PKM2, Unkown	2)GADFLVTEVENGGSLGSK		expressed
protein	3)GVNLPGAAVDLPAVSEK		
	4)TATESFASDPILYRPVAVALDTK		
Sorbitol	1)LENYPIPEPGPNEVLLR	15q15.3cM	Ubiquitously
Dehydrogenase			expressed
SORD			(spermatogenic
			cells)
Lactate	1)DYNVTANSK	11p15.4	Ubiquitous
Dehydrogenase	2)VTLTSEEEAR		
LDHA	3)VIGSGCNLDSAR		
	4)LVIITAGAR		
	5)SADTLWGIQK		
	6)DQLIYNLLKEEQTPQNK		
	7)LKGEMMDLQHGSLFLR		
	8)DLADELALVDVIEDK		
Triosephosphate	1)TATPQQAQEVHEK	12p13	Übiquitously
Isomerase 1	2)LDEREAGITEK		expressed
TPI1	3)IAVAAQNCYK		
	4)SNVSDAVAQSTR		
	5)IIYGGSVTGATCK		
	6)VTNGAFTGEISPGMIK		
	7)HVFGESDELIGQK		
	8)FFVGGNWK		
	9)DCGATWVVLGHSER		
	10)VPADTEVVCAPPTAYIDFAR		
	11)VVLAYEPVWAIGTGK		
	12)QSLGELIGTLNAAK		
	13)KFFVGGNWK		
	14)RHVFGESDELIGQK		
	15)KQSLGELIGTLNAAK		
	16)VAHALAEGLGVIACIGEK		

Figure 9- Expression and Purification of recombinant SFEC

FIG. 9A FIG. 9B FIG. 9C

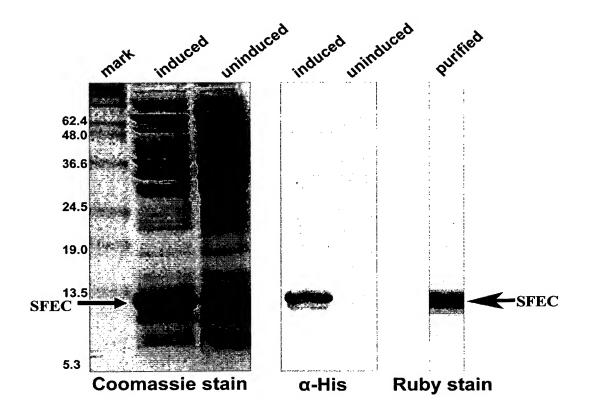


Figure 10- Western Analysis of SFEC

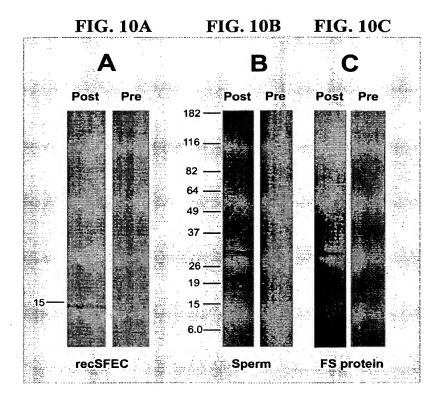


Figure 11- Localization of SFEC to the Principal Piece of the Flagellum

